



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/628,296
Source: OPE
Date Processed by STIC: 8/15/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202



OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/628,296

DATE: 08/05/2003
TIME: 10:08:31

Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

5 <110> APPLICANT: Li, Lian-Chao
7 Cosgrove, Daniel J.
11 <120> TITLE OF INVENTION: PLANT CELL WALL LOOSENING ACTIVITY OF GROUP 2/3 ALLERGENS OF
GRASS POLLEN
15 <130> FILE REFERENCE: P06331US01
C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/628,296
20 <141> CURRENT FILING DATE: 2003-07-28
24 <150> PRIOR APPLICATION NUMBER: US 60,399,688
26 <151> PRIOR FILING DATE: 2002-07-29
30 <160> NUMBER OF SEQ ID NOS: 18
34 <170> SOFTWARE: PatentIn version 3.1
38 <210> SEQ ID NO: 1
40 <211> LENGTH: 291
42 <212> TYPE: DNA
44 <213> ORGANISM: Lolium perenne
48 <400> SEQUENCE: 1
49 acaaaaagtgc atttaactgt ggagaagggt tctgacgcga agacgctgt gctgaacatc 60
51 aagtacacga ggccagggga caccctggcg gaggtggagc tccggcagca cggctcgag 120
53 gagtggaaac ccatgacgaa gaagggcaac ctgtgggagg tgaagagcgc caagccgctc 180
55 accggcccaa tgaacctccg cttcctctcc aaggcggca tgaagaacgt cttcgacgag 240
57 gtcatccccca ccgccttcac ggtcggcaaaa acctacacccc cagaatacaa t 291
60 <210> SEQ ID NO: 2
62 <211> LENGTH: 97
64 <212> TYPE: PRT
66 <213> ORGANISM: Lolium perenne
70 <400> SEQUENCE: 2
72 Thr Lys Val Asp Leu Thr Val Glu Lys Gly Ser Asp Ala Lys Thr Leu
73 1 5 10 15
76 Val Leu Asn Ile Lys Tyr Thr Arg Pro Gly Asp Thr Leu Ala Glu Val
77 20 25 30
80 Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Met Thr Lys Lys
81 35 40 45
84 Gly Asn Leu Trp Glu Val Lys Ser Ala Lys Pro Leu Thr Gly Pro Met
85 50 55 60
88 Asn Phe Arg Phe Leu Ser Lys Gly Gly Met Lys Asn Val Phe Asp Glu
89 65 70 75 80
92 Val Ile Pro Thr Ala Phe Thr Val Gly Lys Thr Tyr Thr Pro Glu Tyr
93 85 90 95
96 Asn
100 <210> SEQ ID NO: 3
102 <211> LENGTH: 291
104 <212> TYPE: DNA
106 <213> ORGANISM: Artificial
W--> 110 <220> FEATURE:
See p. 6 for error explanation

Does Not Comply
Corrected Diskette Needed

ppl-3,6

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Input Set : A:\P06331US01.ST25.txt
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W--> 110 <223> OTHER INFORMATION:

W--> 110 <400> 3

111	acaaaagtgc	atctaactgt	ggagaagggt	tctgacgcga	agacgctgggt	gctgaacatc	60
113	aagtacacga	ggccagggga	caccctggcg	gagggtggagc	tccggcagca	cggtctggag	120
115	gagtgggaac	ccctgacgaa	gaagggcaac	ctgtgggagg	tgaagagcgc	caagccgctc	180
117	accggcccaa	tgaacttccg	cttcctctcc	aaggcgccga	tgaagaacgt	tttcgacgag	240
119	gtcatcccc	ccgccttcac	ggtcggcaaa	acctacaccc	cagaatacaa	t	291

122 <210> SEQ ID NO: 4

124 <211> LENGTH: 97

126 <212> TYPE: PRT

128 <213> ORGANISM: Artificial

W--> 132 <220> FEATURE:

W--> 132 <223> OTHER INFORMATION:

W--> 132 <400> 4

134	Thr Lys Val Asp Leu Thr Val Glu Lys Gly Ser Asp Ala Lys Thr Leu						
135	1	5	10	15			
138	Val Leu Asn Ile Lys Tyr Thr Arg Pro Gly Asp Thr Leu Ala Glu Val						
139		20	25	30			
142	Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Met Thr Lys Lys						
143	35	40	45				
146	Gly Asn Leu Trp Glu Val Lys Ser Ala Lys Pro Leu Thr Gly Pro Met						
147	50	55	60				
150	Asn Phe Arg Phe Leu Ser Lys Gly Gly Met Lys Asn Val Phe Asp Glu						
151	65	70	75	80			
154	Val Ile Ala Thr Ala Phe Thr Val Gly Lys Thr Tyr Thr Pro Glu Tyr						
155	85	90	95				

158 Asn

162 <210> SEQ ID NO: 5

164 <211> LENGTH: 20

166 <212> TYPE: PRT

168 <213> ORGANISM: Lolium perenne

172 <400> SEQUENCE: 5

174	Thr Lys Val Asp Leu Thr Val Glu Lys Gly Ser Asp Ala Lys Thr Leu						
175	1	5	10	15			

178 Val Leu Asn Ile

179 20

182 <210> SEQ ID NO: 6

184 <211> LENGTH: 52

186 <212> TYPE: DNA

188 <213> ORGANISM: Artificial

192 <220> FEATURE:

194 <221> NAME/KEY: misc_feature

196 <222> LOCATION: (3)..(3)

198 <223> OTHER INFORMATION: N is unknown

202 <220> FEATURE:

204 <221> NAME/KEY: misc_feature

206 <222> LOCATION: (15)..(15)

208 <223> OTHER INFORMATION: N is unknown

212 <220> FEATURE:

See p. 6

Needs explanation - see p. 6

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Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

214 <221> NAME/KEY: misc_feature
 216 <222> LOCATION: (29)..(29)
 218 <223> OTHER INFORMATION: N is unknown
 222 <220> FEATURE:
 224 <221> NAME/KEY: misc_feature
 226 <222> LOCATION: (36)..(36)
 228 <223> OTHER INFORMATION: N is unknown
 232 <220> FEATURE:
 234 <221> NAME/KEY: misc_feature
 236 <222> LOCATION: (43)..(43)
 238 <223> OTHER INFORMATION: N is unknown
 242 <400> SEQUENCE: 6
 W--> 243 acnacgtaar aggtacgtg ayctycttna cgtacnacgt gtnacgtgar ag 52
 246 <210> SEQ ID NO: 7
 248 <211> LENGTH: 49
 250 <212> TYPE: DNA
 252 <213> ORGANISM: Artificial needs explanation - see p. 6
 256 <220> FEATURE:
 258 <221> NAME/KEY: misc_feature
 260 <222> LOCATION: (24)..(24)
 262 <223> OTHER INFORMATION: N is unknown
 266 <220> FEATURE:
 268 <221> NAME/KEY: misc_feature
 270 <222> LOCATION: (31)..(31)
 272 <223> OTHER INFORMATION: N is unknown
 276 <220> FEATURE:
 278 <221> NAME/KEY: misc_feature
 280 <222> LOCATION: (43)..(43)
 282 <223> OTHER INFORMATION: N is unknown
 286 <400> SEQUENCE: 7
 W--> 287 cyctyctara gttragtayc ttcnacgtgg nacgtgtrag tanacgtgt 49
 290 <210> SEQ ID NO: 8
 292 <211> LENGTH: 93
 294 <212> TYPE: PRT
 296 <213> ORGANISM: Zea mays
 300 <400> SEQUENCE: 8
 302 Thr Phe Gln Val Gly Lys Ser Lys Pro Gly His Leu Val Leu Thr
 303 1 5 10 15
 306 Pro Asn Ile Ala Thr Ile Ser Asp Val Glu Ile Lys Glu His Gly Gly
 307 20 25 30
 310 Asp Asp Phe Ser Phe Thr Leu Lys Glu Gly Pro Ala Gly Thr Trp Thr
 311 35 40 45
 314 Leu Asp Thr Lys Ala Pro Leu Lys Tyr Pro Leu Cys Ile Arg Phe Ala
 315 50 55 60
 318 Thr Lys Ser Gly Gly Tyr Arg Ile Ala Asp Asp Val Ile Pro Ala Asp
 319 65 70 75 80
 322 Phe Lys Ala Gly Thr Thr Tyr Lys Thr Thr Leu Ser Ile
 323 85 90
 326 <210> SEQ ID NO: 9

Please ensure all 213 responses of Artificial or of Unknown are explained in 220-223 section

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Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

328 <211> LENGTH: 96
 330 <212> TYPE: PRT
 332 <213> ORGANISM: Dactylis glomerata
 336 <400> SEQUENCE: 9
 338 Val Lys Val Thr Phe Lys Val Glu Lys Gly Ser Asp Pro Lys Lys Leu
 339 1 5 10 15
 342 Val Leu Asp Ile Lys Tyr Thr Arg Pro Gly Asp Thr Leu Ala Glu Val
 343 20 25 30
 346 Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Leu Thr Lys Lys
 347 35 40 45
 350 Gly Asn Leu Trp Glu Val Lys Ser Ser Lys Pro Leu Thr Gly Pro Phe
 351 50 55 60
 354 Asn Phe Arg Phe Met Ser Lys Gly Gly Met Arg Asn Val Phe Asp Glu
 355 65 70 75 80
 358 Val Ile Pro Thr Ala Phe Lys Ile Gly Thr Thr Tyr Thr Pro Glu Glu
 359 85 90 95
 362 <210> SEQ ID NO: 10
 364 <211> LENGTH: 90
 366 <212> TYPE: PRT
 368 <213> ORGANISM: Oryza sativa
 372 <400> SEQUENCE: 10
 374 Met Glu Val Ala Lys Gly Ser Ser Ala Lys Ser Leu Glu Leu Val Thr
 375 1 5 10 15
 378 Asn Val Ala Ile Ser Lys Val Glu Val Lys Glu Lys Gly Lys Asp
 379 20 25 30
 382 Trp Val Ala Leu Lys Glu Ser Ser Asn Thr Trp Thr Leu Lys Ser
 383 35 40 45
 386 Glu Ser Pro Leu Lys Gly Pro Phe Ser Val Arg Phe Leu Val Lys Asn
 387 50 55 60
 390 Ser Gly Tyr Arg Val Val Asp Asp Ile Ile Pro Glu Ser Phe Thr Ala
 391 65 70 75 80
 394 Gly Ser Glu Tyr Lys Ser Gly Ile Gln Leu
 395 85 90
 398 <210> SEQ ID NO: 11
 400 <211> LENGTH: 96
 402 <212> TYPE: PRT
 404 <213> ORGANISM: Lolium perenne
 408 <400> SEQUENCE: 11
 410 Ala Ala Pro Val Glu Phe Thr Val Glu Lys Gly Ser Asp Glu Lys Asn
 411 1 5 10 15
 414 Leu Ala Leu Ser Ile Lys Tyr Asn Lys Glu Gly Asp Ser Met Ala Glu
 415 20 25 30
 418 Val Glu Leu Lys Glu His Gly Ser Asn Glu Trp Leu Ala Leu Lys Lys
 419 35 40 45
 422 Asn Gly Asp Gly Val Trp Glu Ile Lys Ser Asp Lys Pro Leu Lys Gly
 423 50 55 60
 426 Pro Phe Asn Phe Arg Phe Val Ser Glu Lys Gly Met Arg Asn Val Phe
 427 65 70 75 80
 430 Asp Asp Val Val Pro Ala Asp Phe Lys Val Gly Thr Thr Tyr Lys Pro

RAW SEQUENCE LISTING
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Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

431	85	90	95
434 <210> SEQ ID NO: 12			
436 <211> LENGTH: 96			
438 <212> TYPE: PRT			
440 <213> ORGANISM: Sorghum			
444 <400> SEQUENCE: 12			
446 Gly Thr Thr Leu Thr Ile Glu Val Gly Lys Asp Ser Thr Ser Thr Lys			
447 1 5 10 15			
450 Leu Ser Leu Ile Thr Asn Val Ala Ile Ser Glu Val Ser Val Lys Pro			
451 20 25 30			
454 Lys Gly Ala Thr Asp Phe Thr Asp Asp Leu Lys Glu Ser Glu Pro Lys			
455 35 40 45			
458 Thr Phe Thr Leu Asp Ser Lys Glu Pro Ile Glu Gly Pro Ile Ala Phe			
459 50 55 60			
462 Arg Phe Leu Ala Lys Gly Gly Tyr Arg Val Val Asp Asn Ala Ile			
463 65 70 75 80			
466 Pro Ala Asp Phe Lys Ala Gly Ser Val Tyr Lys Thr Thr Glu Gln Val			
467 85 90 95			
470 <210> SEQ ID NO: 13			
472 <211> LENGTH: 100			
474 <212> TYPE: PRT			
476 <213> ORGANISM: Hordeum vulgare			
480 <400> SEQUENCE: 13			
482 Ala Ala Thr Lys Val Lys Phe Thr Val Gln Lys Gly Ser Asp Ala Lys			
483 1 5 10 15			
486 Lys Leu Val Leu Lys Ile Asp Tyr Thr Arg Ala Gly Asp Thr Leu Ser			
487 20 25 30			
490 Glu Met Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Phe Thr			
491 35 40 45			
494 Lys Lys Gly Asp Val Trp Glu Leu Ser Ser Lys Pro Leu Val Gly			
495 50 55 60			
498 Pro Phe Asn Phe Arg Phe Leu Ser Lys Gly Gly Met Lys Asn Val Phe			
499 65 70 75 80			
502 Asp Glu Val Phe Ser Thr Asp Phe Lys Ile Gly Lys Thr Tyr Glu Pro			
503 85 90 95			
506 Val Tyr Asp Ala			
507 100			
510 <210> SEQ ID NO: 14			
512 <211> LENGTH: 98			
514 <212> TYPE: PRT			
516 <213> ORGANISM: Triticum aestivum			
520 <400> SEQUENCE: 14			
522 Ala Pro Pro Pro Val Ser Ile Thr Val Glu Lys Gly Ser Asp Ala Lys			
523 1 5 10 15			
526 His Leu Val Leu Gln Ile Lys Tyr Asp Lys Val Gly Asp Ser Met Lys			
527 20 25 30			
530 Glu Val Glu Leu Glu Gln Asn Glu Asp Trp Leu Pro Leu Lys Lys Gly			
531 35 40 45			
534 Tyr Ser Gly Ala Trp Glu Ile Lys Ser Asp Thr Pro Leu Lys Gly Pro			

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/05/2003
PATENT APPLICATION: US/10/628,296 TIME: 10:08:32

Input Set : A:\P06331US01.ST25.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 3,15,29,36,43
Seq#:7; N Pos. 24,31,43
Seq#:17; Xaa Pos. 27

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 11

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,6,7

Use of <220> Feature(NEW RULES): *error explanation*
Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:3,4,6,7


VERIFICATION SUMMARY
PATENT APPLICATION: US/10/628,296

DATE: 08/05/2003
TIME: 10:08:32

Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

L:19 M:270 C: Current Application Number differs, Wrong Format
L:110 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:3, <213>
ORGANISM:Artificial
L:110 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:3, <213>
ORGANISM:Artificial
L:110 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:110
L:132 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:4, <213>
ORGANISM:Artificial
L:132 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:4, <213>
ORGANISM:Artificial
L:132 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:132
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:652 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:16